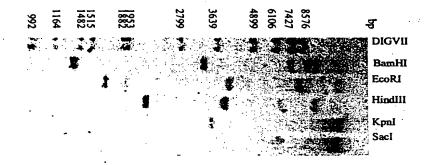


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Soybean genomic DNA Southern blot hybridized with the SAMS promoter probe.

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7427 6106 4899 3639 2799 11515 1164

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DIGVII

BamHI

EcoRI

HindIII

Kpnl

SacI

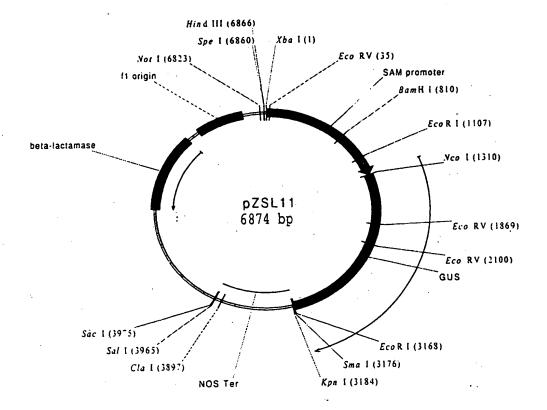
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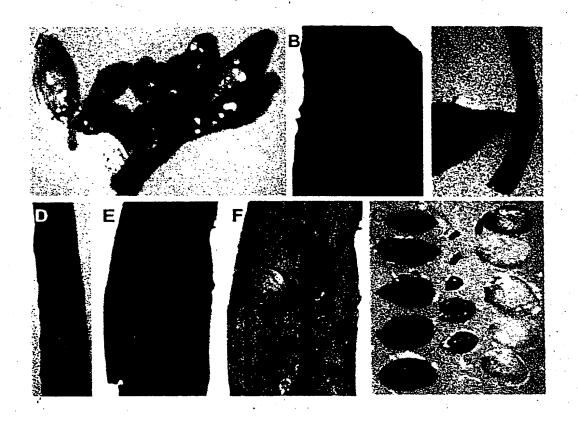
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AGATGCAGAGACATTCCTATTCACCTCGGAGTCAGTGAACGAGGGACACCCTGATAAGCTCTGCGACCAAATCTCCGAT

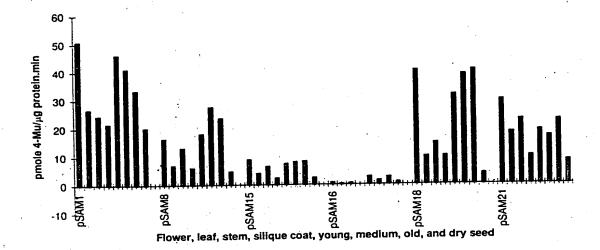
 ${\tt GCTGTCCTCGACGCTTGCCTCGAACAGGACCCAGACAGCAAGGTTGCCTGCGAAACATGCACCAAGACCAACTTGGTCAT}$ GCTGTCCTCGACGCTTGCCCTTGAACAGGACCCAGACAGCAAGGTTGCCTGCGAAACATGCACCAAGACCAACTTGGTCAT

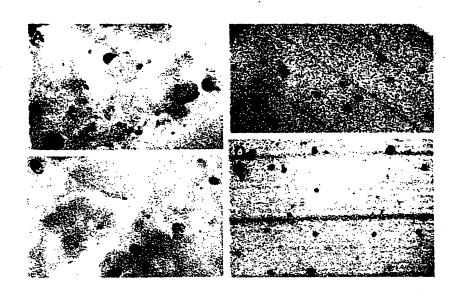
GGTCTTCGGAGAGATC GGTCTTCGGAGAGATC

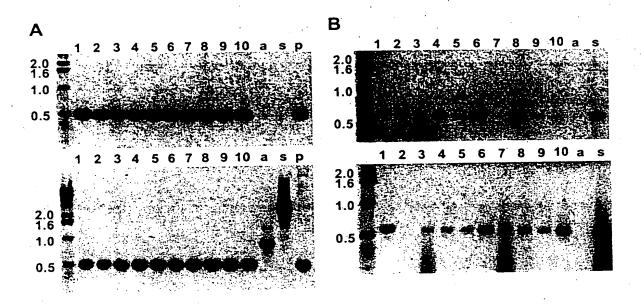












0.01 mM, and 1 mM methionine. Ten days old seedlings were harvested and assayed for GUS activity. The solid with SAMS::GUS construct were germinated on filter papers soaked with H₂O, 1x Murashige and Skoog salt,

Induction of SAMS promoter activity by free methionine. Seeds of ten transgenic Arabidopsis lines transformed

bar and hollow bar indicate, respectively, the average and the standard variation of three samples for treatment.

